

--6. (New) The microorganism of claim 1 wherein the microorganism belonging to enterobacteria is a bacterium belonging to the genus *Enterobacter* or *Klebsiella*.

7. (New) The microorganism of claim 2 wherein the microorganism belonging to enterobacteria is a bacterium belonging to the genus *Enterobacter* or *Klebsiella*.

8. (New) The microorganism of claim 6 wherein the bacterium is *Enterobacter agglomerans* or *Klebsiella planticola*.

9. (New) The microorganism of claim 7 wherein the bacterium is *Enterobacter agglomerans* or *Klebsiella planticola*.

10. (New) A process for producing L-glutamic acid comprising the steps of culturing the microorganism of claim 1 in a liquid medium to produce and accumulate L-glutamic acid in the medium and collecting the L-glutamic acid from the medium.

*Sub C2*

11. (New) A process for producing L-glutamic acid comprising isolating a coryneform bacterium citrate synthase gene by amplifying the gene with oligonucleotide primers comprising SEQ ID NOS: 1 and 2; transforming a enterobacteria with said isolated citrate synthase gene; culturing said enterobacteria in a liquid medium to produce and accumulate the L-glutamic acid; and collecting the L-glutamic acid produced.

12. (New) The process of Claim 11, wherein the coryneform bacteria is *Brevibacterium lactofermentum*.

13. (New) The process of Claim 11, wherein the entereobacteria is of the genus *Enterobacter* or *Klebsiella*.

14. (New) The process of Claim 11, wherein the enterobacteria is *Enterobacter agglomerans* or *Klebsiella planticola*.

*Add C3*